

- 1/2 -

Fig. 1**SEQ ID NO:1**

5' - ACTGTTCTGAGATCTTTGCCTCCCTCAGGCTCCCGAGAATCATGGCTCAT
TCCAAGCAGCCTAGTCACTTCCAGAGTCTGATGCTTCTGCAGTGGCCTTT
GAGCTACCTTGCCATCTTTTGGATCTTGCAGCCATTGTTTCGTCTACCTGC
TGTTTACATCCTTGTGGCCGCTACCAGTGCTTTACTTTGCCTGGTTGTTT
CTGGACTGGAAGACCCAGAGCGAGGTGGCAGGCGTTCGGCCTGGGTAAG
GAACTGGTGTGTCTGGACCCACATCAGGGACTATTTCCCCATTACGATCC
TGAAGACAAAGGACCTATCACCTGAGCACAACTACCTCATGGGGGTTCAC
CCCCATGGCCTCCTGACCTTTGGCGCCTTCTGCAACTTCTGCACTGAGGC
CACAGGCTTCTCGAAGACCTTCCCAGGCATCACTCCTCACTTGGCCACGC
TGTCTGGTTCTTCAAGATCCCCTTTGTTAGGGAGTACCTCATGGCCAAA
GGTGTGTGCTCTGTGAGCCAGCCAGCCATCAACTATCTGCTGAGCCATGG
CACTGGCAACCTCGTGGGCATTGTAGTGGGAGGTGTGGGTGAGGCCCTGC
AAAGTGTGCCCAACACCACCACCCTCATCCTCCAGAAGCGCAAGGGGTTC
GTGCGCACAGCCCTCCAGCATGGGGCTCATCTGGTCCCCACCTTCACTTT
TGGGGAAACTGAGGTGTATGATCAGGTGCTGTTCCATAAGGATAGCAGGA
TGTACAAGTTCCAGAGCTGCTTCCGCCGTATCTTTGGTTTCTACTGTTGT
GTCTTCTATGGACAAAGCTTCTGTCAAGGCTCCACTGGGCTCCTGCCATA
CTCCAGGCCTATTGTCACTGTGGTTGGGGAGCCTCTGCCACTGCCCCAAA
TTGAAAAGCCAAGCCAGGAGATGGTGGACAAATACCATGCACTTTATATG
GATGCTCTGCACAAACTGTTTCGACCAGCATAAGACCCACTATGGCTGCTC
AGAGACCCAAAAGCTGTTTTTCCTGTGAATGAAGGTACTGCATGCCCAGG
AGCACAGGAGTGCCTGCCTTGAAGAAGAGACTCATCTGCCACTAACCAAA
GACAGGCAGGAGATGAGGGAGGTTATATGTG

3'

- 2/2 -

Fig. 2:**SEQ ID NO:2**

MAHSKQPSHFQSLMLLQWPLSYLAIFWILQPLFVYLLFTSLWPLPVLYFA
WLFLDWKTPERGGRRSAWVRNWCVWTHIRDYFPITLKT KDLSPEHNYLM
GVHPHGLLTFGAFCNFCTEATGFSKTFPGITPHLATLSWFFKIPFVREYL
MAKGVCSVSQPAINYLLSHGTGNLVGIVVGGVGEALQSV PNTTTLILQKR
KGFVRTALQHGAHLVPTFTFGETEVYDQVLFHKDSRMYK FQSCFRRIFGF
YCCVFY GQSFCQGSTGLLPYSRPIVTVVGEPLPLPQIEKPS QEMVDKYHA
LYMDALHKLFDQHKTHYGCSETQKLFFL

Fig. 3:**SEQ ID NO:3**

5'-CCAGGCCTATTGTC ACTGTG-3'

Fig. 4:**SEQ ID NO:4**

5'-CTGGCTTGGCTTTTCAATTT-3'

Fig. 5:**SEQ ID NO:5**

5'-TTGGGGAGCCTCTGCCACTGC-3'